



Supplementary information, Fig. S3 Representative cryo-EM densities of the cGAS-nucleosome complex. **a-d** Cryo-EM densities of H2A helix (**a**), H2B helix (**b**), cGAS Loop 1 (**c**) and cGAS Loop 2 (**d**). The cryo-EM densities were generated from the 3.8 Å map of the overall monomer. **e** Cryo-EM density of cGAS-CD. The cryo-EM density was generated from the 4.7 Å map of the subtracted monomer.